

FIGURE 1A

CTCGAGGACAGTGACCTGGGAGTGAGTACAAGGTGAGGCCACCACTCAGGGT
GCCAGCTCCAAGCGGGTCACAGGGACGAGGGCTGCGGCCATCAGGAGGCCCT
GCACACACATCTGGGACACGCGCCCCGAGGGCCAGTTCACCTCAGTGCGCCT
CATTCTCCTGCACAAAAGCGCCCCATCCTTCTTCACAAGGCTTCGTGGAAG
CAGAGGCCTCGATGCCAGTACCCCTCCCTTCCCAGGCAACGGGACCCCAA
GTTTGCTGACTGGGACCAAGCCACGCATGCGTCAAGAGTGAGAGTCCGG
GACCTAGGCAGGGGCCCTGGGGTTGGGCTGAGAGAGAAGAGAACCTCCCC
AGCACTCGGTGTGCATCGGTAGTGAAGGAGCCTCACCTGACCCCCGCTGTTGC
TCAATCGACTTCCAAGAACAGAGAGAAAAGGAACTTCCAGGGCGGCCGG
GCCTCCTGGGGTTCCCACCCATTCTAGCTGAAAGCACTGAGGCAGAGCTC
CCCCTACCCAGGCTCCACTGCCGGCACAGAAATAACAACCACGGTTACTGAT
CATCTGGGAGCTGTCCAGGAATT

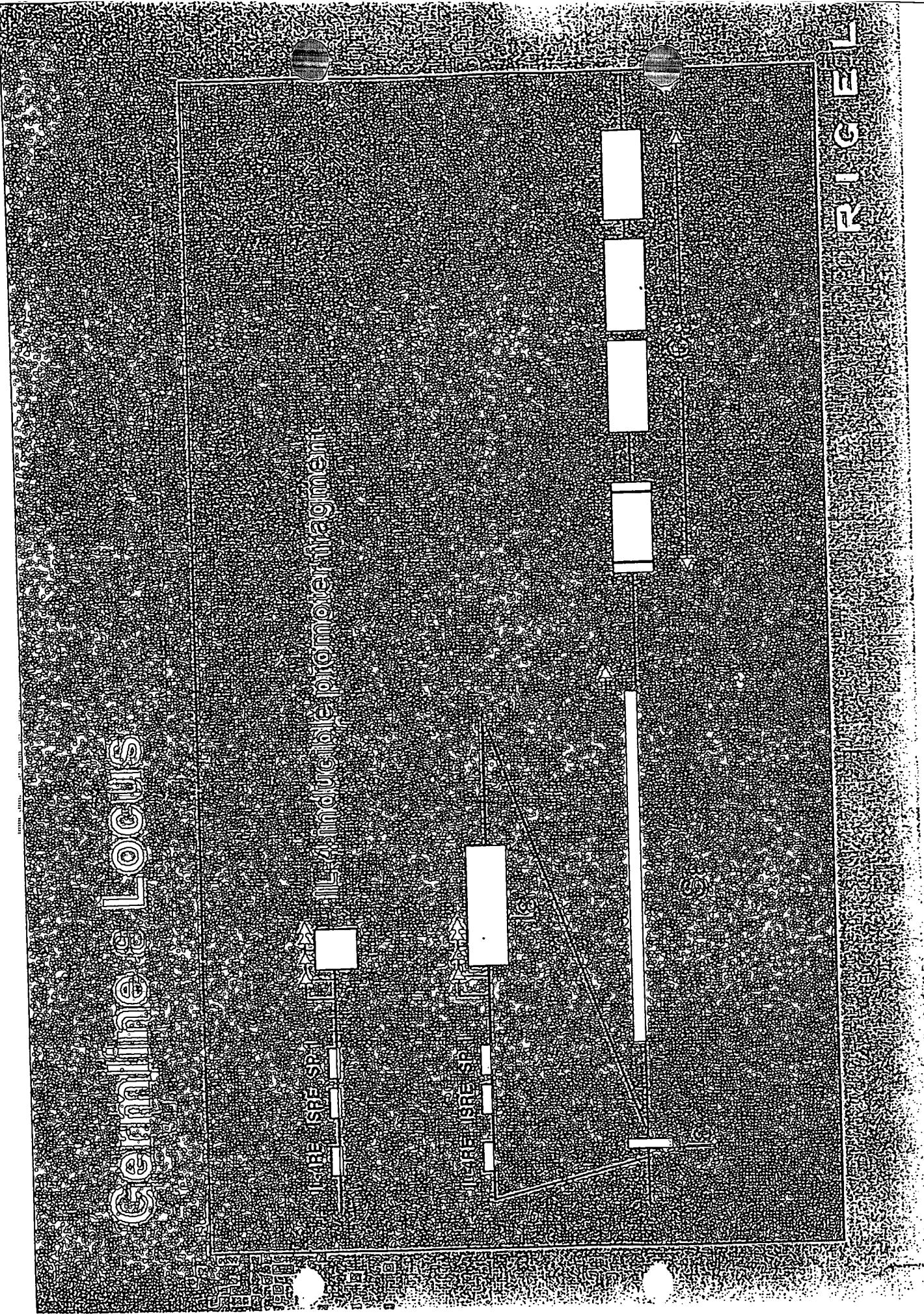


FIGURE 1B

Low energy DNA folding of the S_c region

FIG 2A

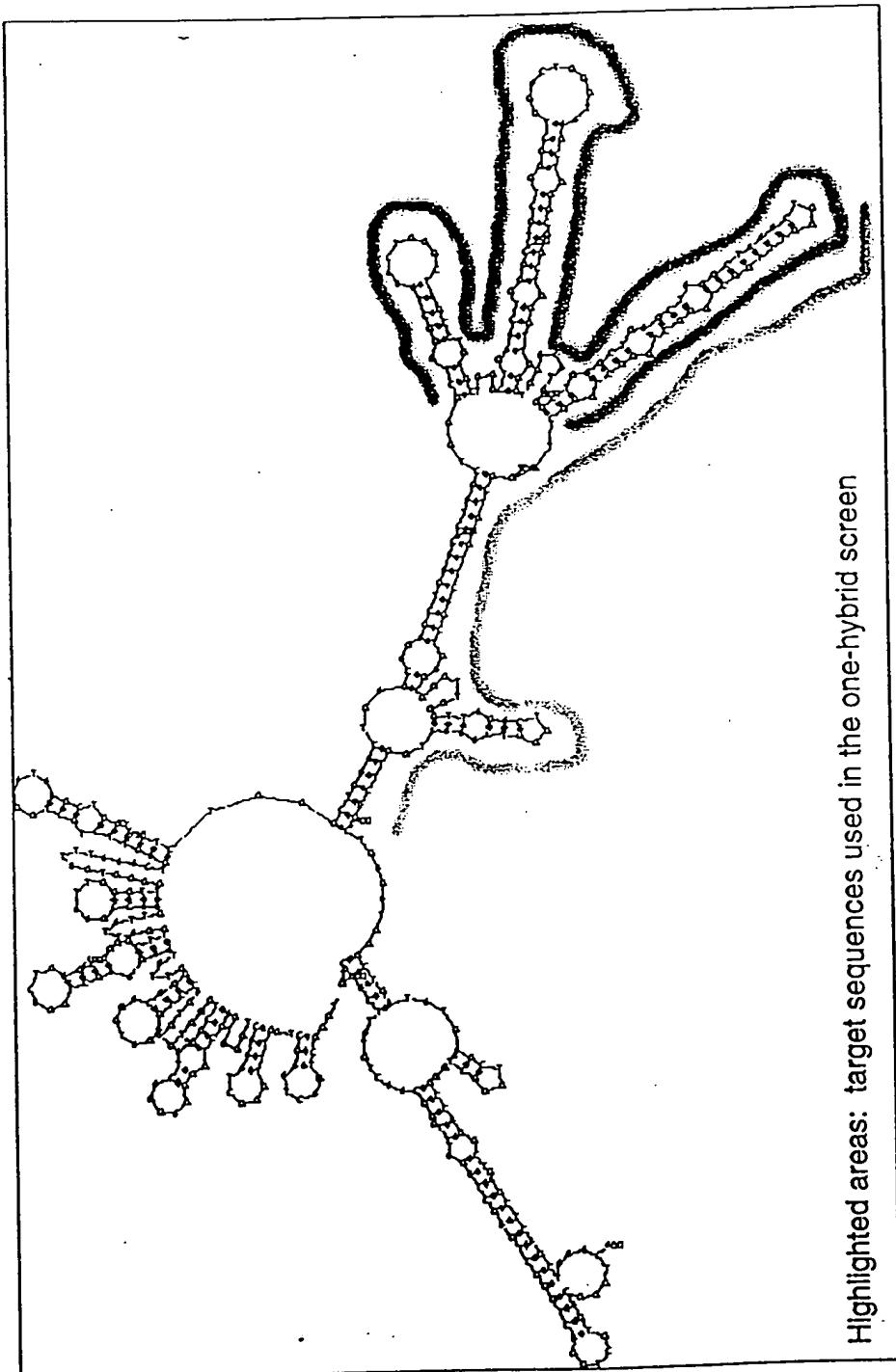


FIGURE 2B

1 GCTGGGCTAA ACTGGGCTAG CCTGAGCTGG GCTGAACCTGG GCTGCTGGC
51 TGGACTGGGT AAGCTGGCT GAGCTGGTT GGGTGGAAAT GGGCTGAGCT
101 GAGCTAGGCT AACTGGGTT TGGCTGGCT GGGCTGGC GGG

FIGURE 2C

1 GGTTGGCTG GGCTGGCTG GGCTGGCTG GGTCAGCTG AGCGGGTTGG
51 GTTAGACTGG GTCAAACTGG TTCAGC

FIG 3

Appendix F

Yeast One-Hybrid Screening

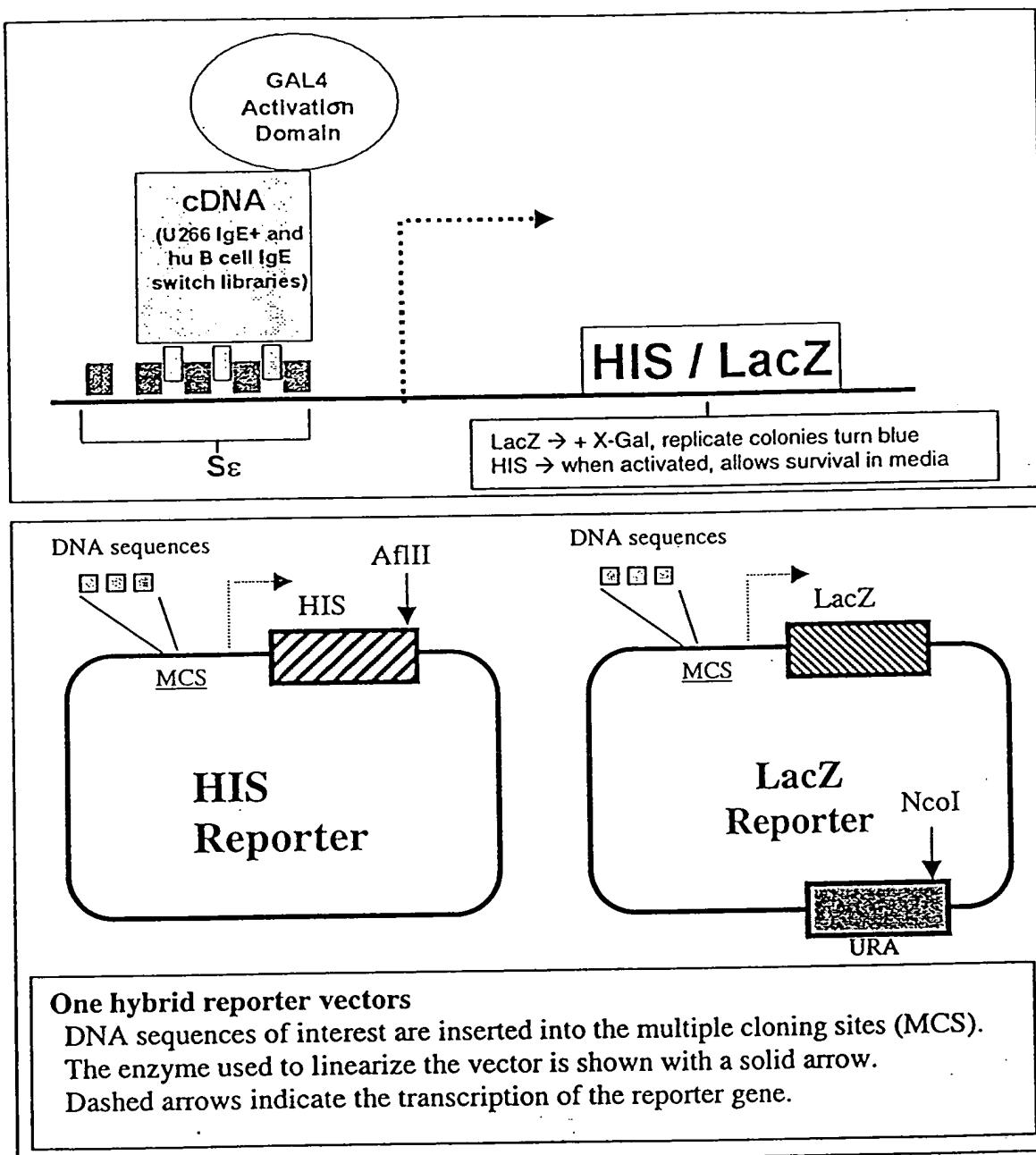
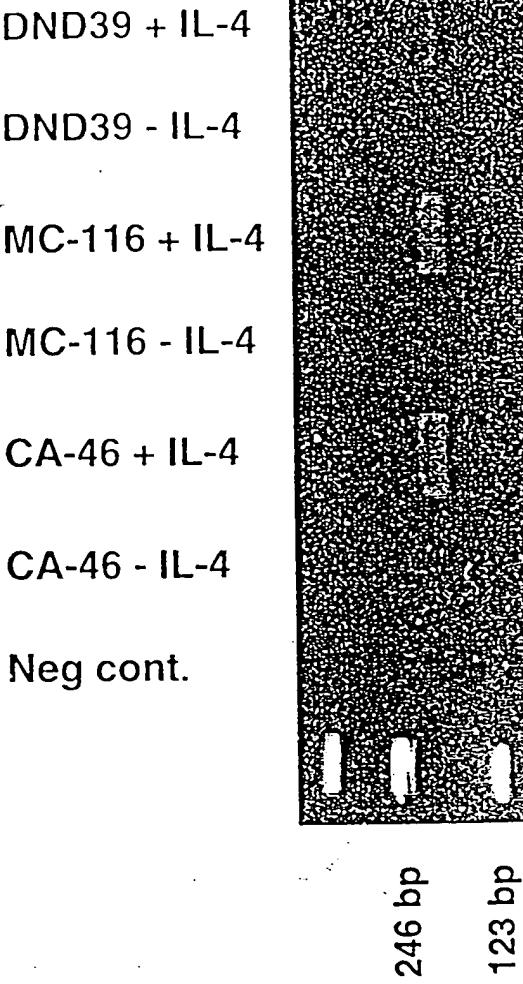


FIG 4

IL-4 Induction of Germline ϵ mRNA in the IgM+ B cell lines: CA-46, MC-116 and DND39

Cells were incubated for 48 hrs in 300 U/ml of h-IL-4. RT-PCR was performed using primers specific for the germline ϵ exon and the 5'-end of the ϵ CH1 exon (predicted size ~ 200 bp).



Approaches to generate germline ϵ promoter knock-in reporter cell lines

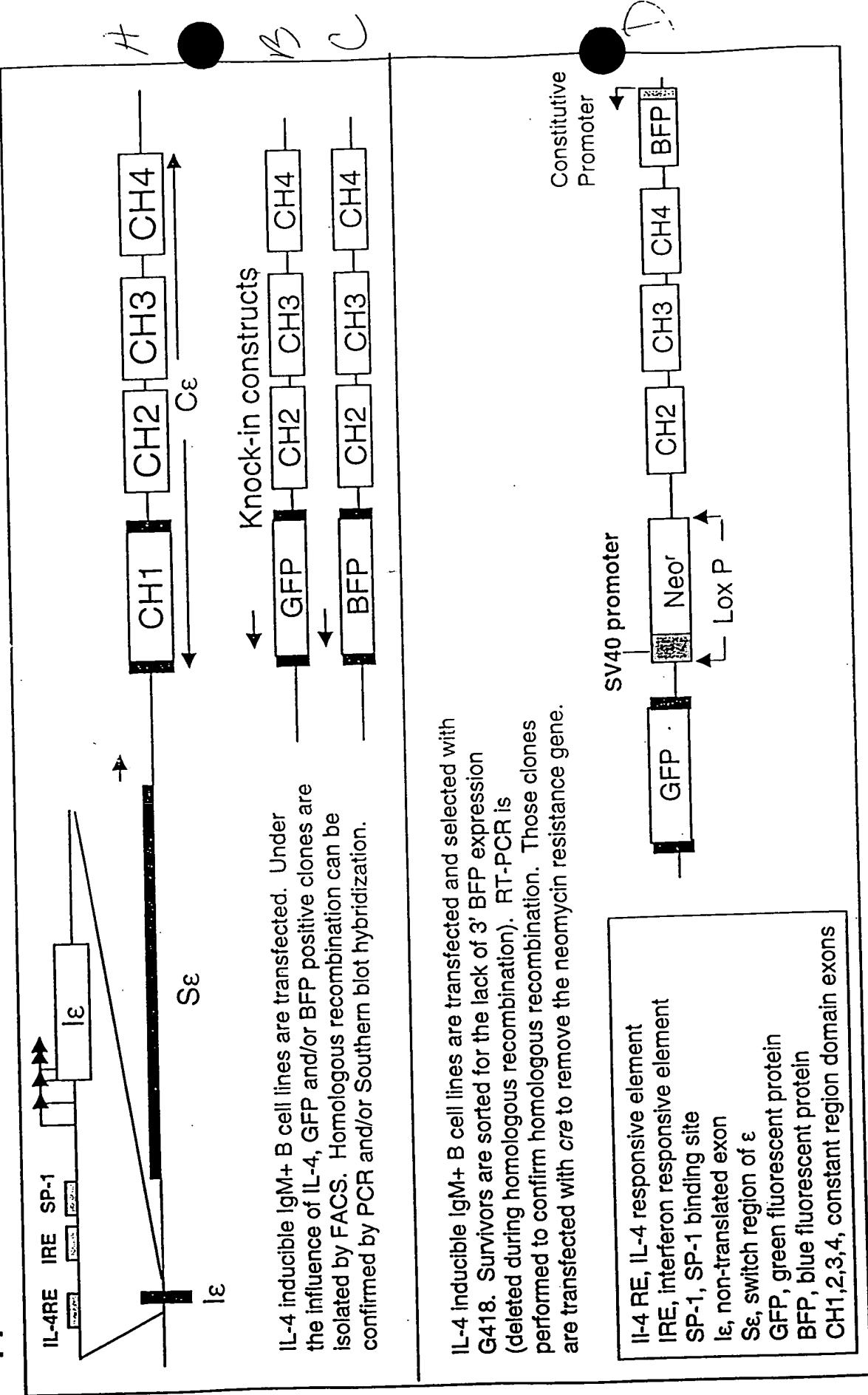
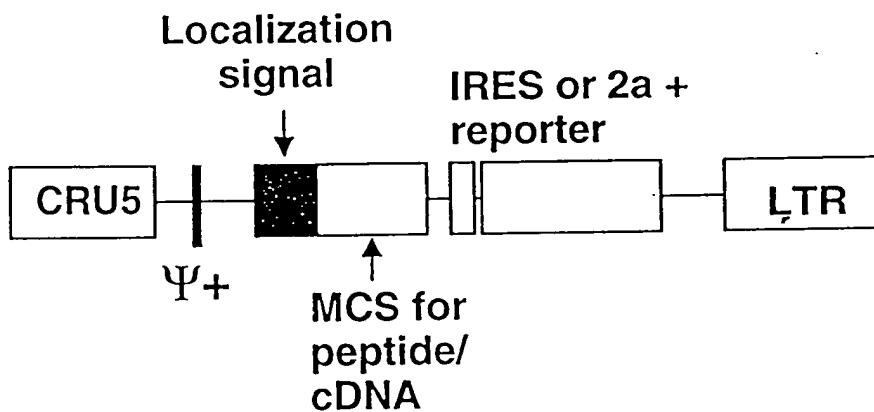


FIG 6

Appendix I

Rigel Base Vector



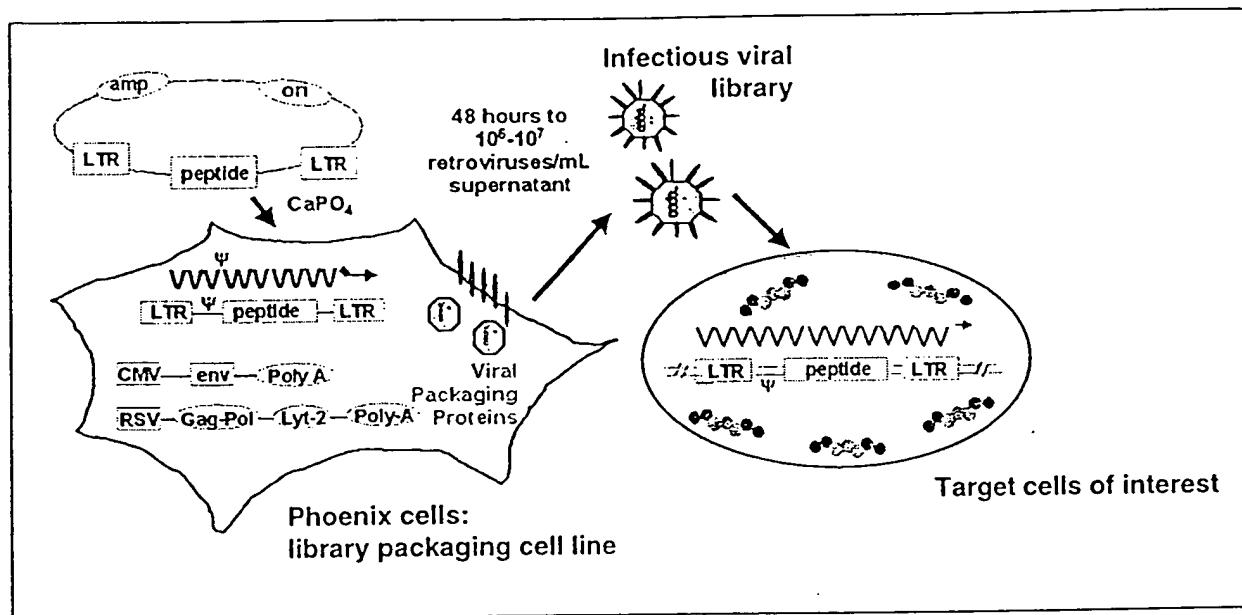
All components are cassetted for flexibility

CRU5, modified LTR
LTR, long terminal repeat
 $\psi+$, packaging signal
Localization signal: nuclear, cell membrane, granular
MCS, multiple cloning site
IRES, internal ribosome entry site
2a, self-cleaving peptide

FIG 7

Appendix H

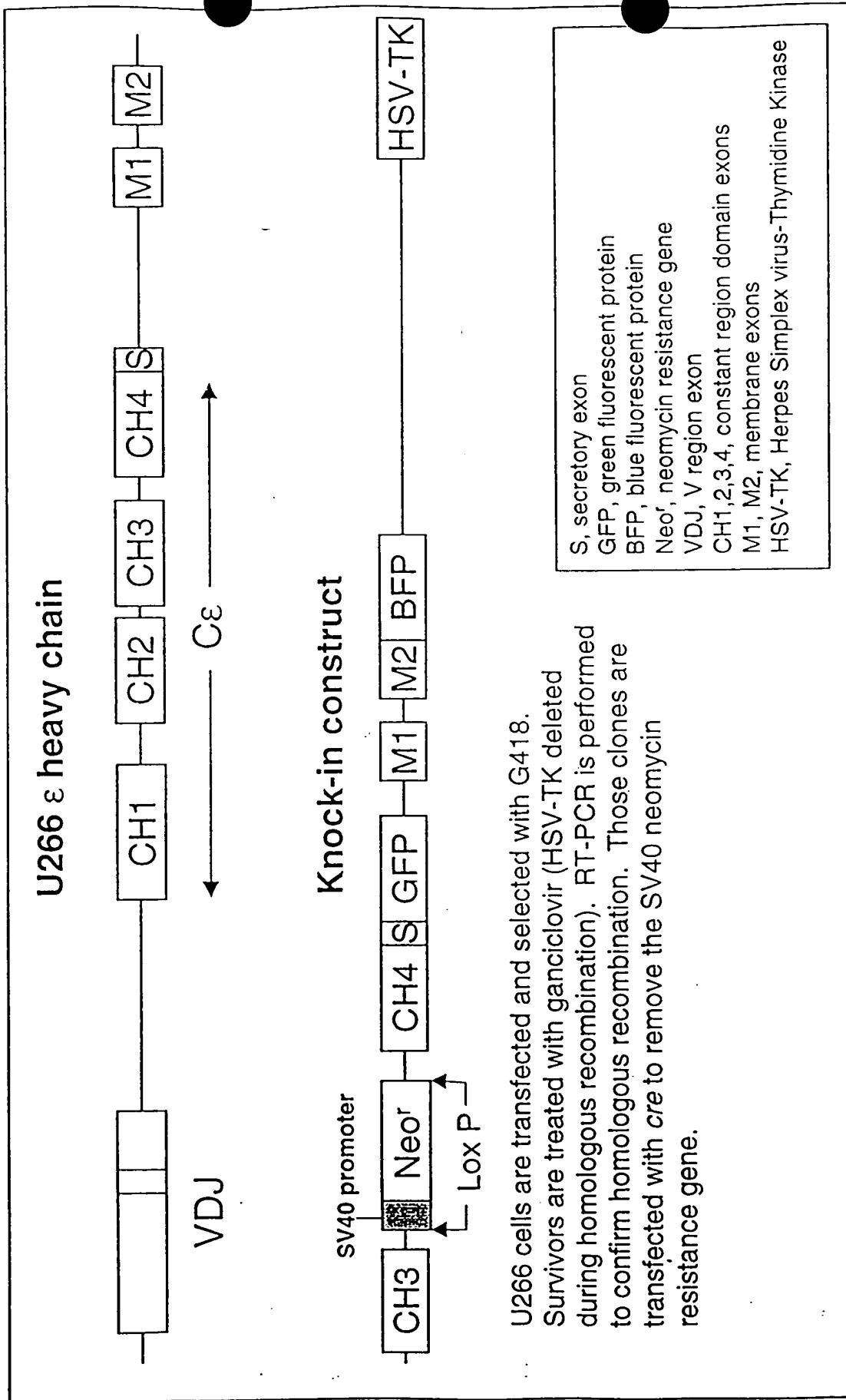
Protocol for Transfection of Phoenix Cells and Infection of Nonadherent Target Cells



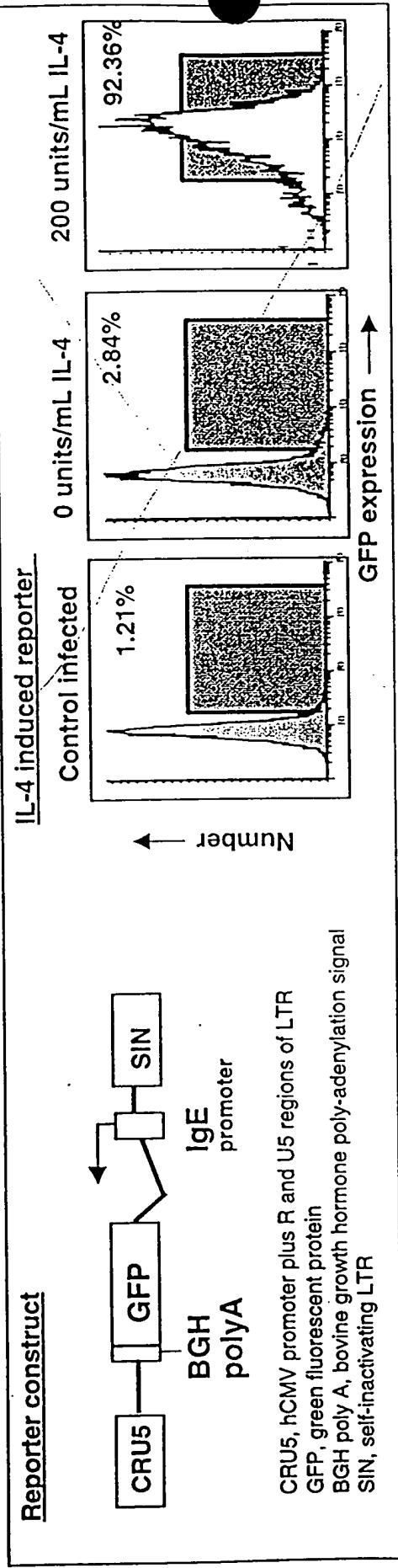
F16 8

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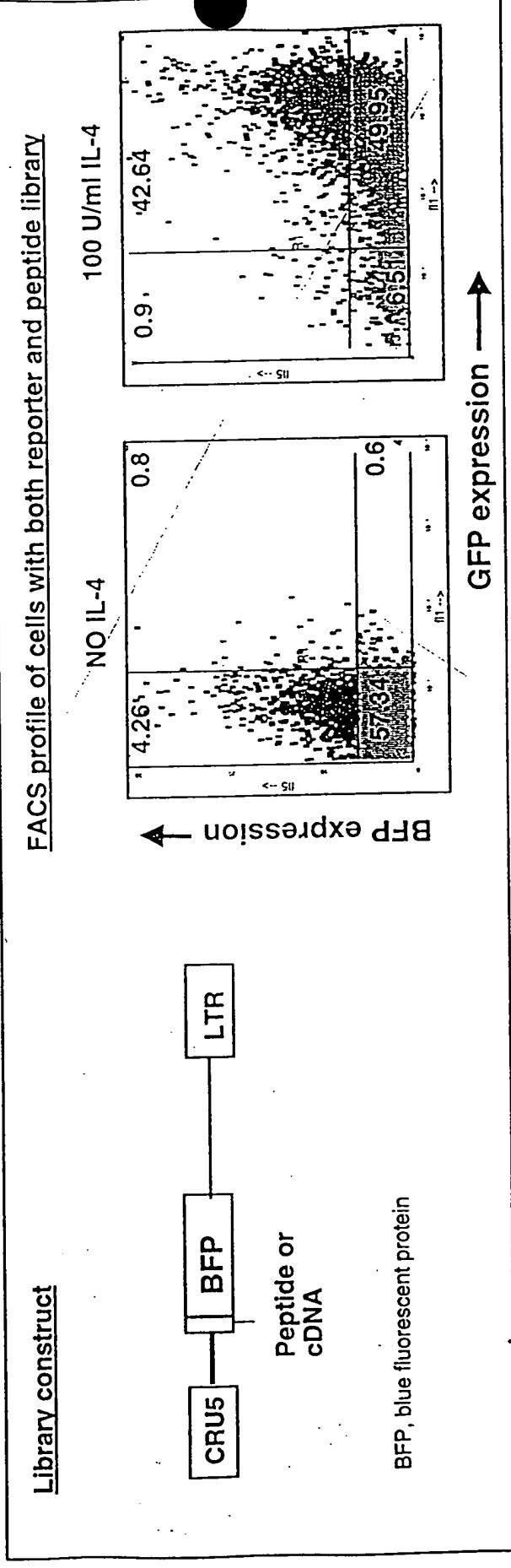
ϵ heavy chain GFP/BFP knock-in cell line



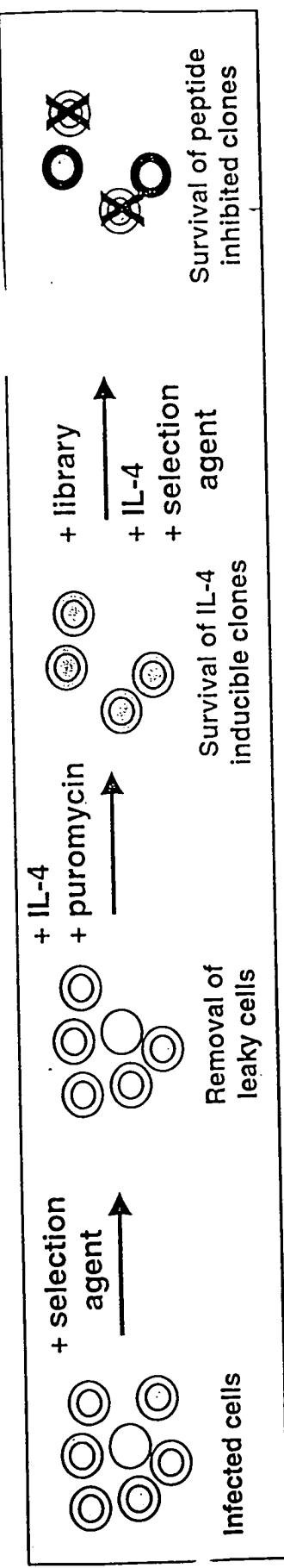
IL-4 Inducible & Promoter Reporter Cell Line



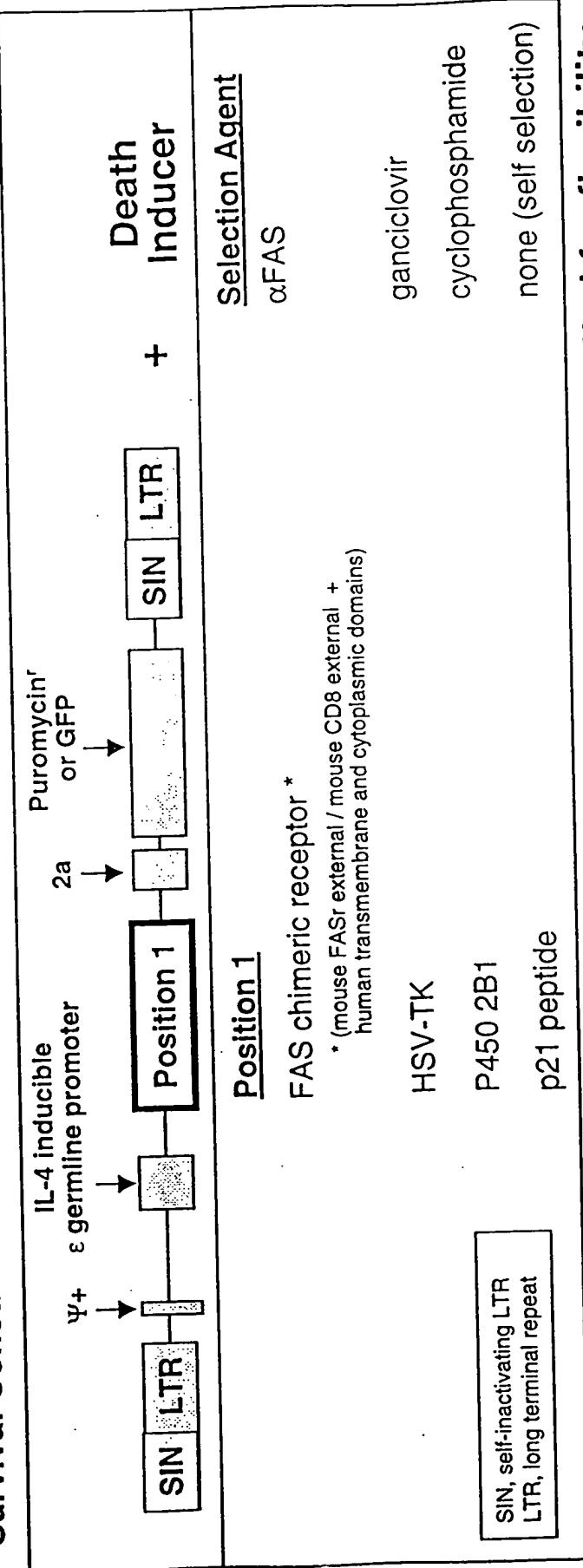
Reporter Line Infected with BFP Construct



Screen for Peptide Inhibitors of the Germline ϵ Promoter



Survival Construct



All components are cassetted for flexibility

Appendix D

FIGURE 11A-1

1-845 CMV promoter/R/U5 5' LTR
1322 GAG ATG-ATC mutation
850-2100 extended ψ region
2146-2173 two BstX1 peptide cloning sites
2205-2723 ECMV IRES (cloned as EcoR1/Msc1 fragment from
pCITE-4a [Novagen])
2746-3465 GFP coding region
3522-4115 3' LTR
4122-6210 pGEM backbone (pUC origin, ampr)

ATCACGAGGCCCTTCGTCTCAAGAACAGCTTGCTCTAGGAGTTCTAATACATCC
CAAACCTAAATATATAAAGCATTGACTTGTCTATGCCCTAGTTATTAAATAGTAATCAA
TTACGGGGTCATTAGTCATAGCCCATATATGGAGTTCCCGCTTACATAACTTACGGTAA
ATGGCCCGCCTGGCTGACGCCAACGACCCCCGCCATTGACGTCAATAATGACGTATG
TTCCCATAGTAACGCAAATAGGGACTTCCATTGACGTCAATGGGTGGAGTATTACGGT
AAACTGCCACTGGCAGTACATCAAGTGTATCATATGCCAAGTACGCCCTATTGACG
TCAATGACGGTAAATGGCCCGCCTGGCATTATGCCAGTACATGACCTTATGGGACTTC
CTACTGGCAGTACATCTACGTATTAGTCATCGTATTACCATGGTATGCCGTTTGGC
AGTACATCAATGGCGTGGATAGCGGTTGACTCACGGGATTCCAAGTCTCCACCCCA
TTGACGTCAATGGGAGTTGTTGGCACCAAAATCAACGGGACTTCCAAAATGTCGTA
ACAACCTCCGCCCATGACGCAAATGGCGGTAGGCATGTACGGTGGGAGGTCTATATAA
GCAGAGCTCAATAAAAGAGCCCACAACCCCTCACTGGGGGCCAGTCTCCGATTGACT
GAGTCGCCGGTACCGTGTATCCAATAAAACCCCTTGCAGTTGCATCCGACTTGTGGT
CTCGCTGTTCTGGGAGGGTCTCTGAGTGTACTACCCGTCAAGGGGGACCCGACCCACCG
CATTTGGGGGCTCGTCCGGATCGGAGACCCCTGCCAGGGACCCGACCCACCG
GGAGGTAAGCTGCCAGCAACTTATCTGTTGCTCGATTTGACTAGTGTCTATGACTGA
TTTATGCGCTGCGTGGTACTAGTTAGCTAACTAGCTCTGTATCTGGCGGACCCGTGG
TGGAACTGACGAGTTCGGAACACCCGGCCCAACCTGGGAGACGTCCCAGGGACTTCGG
GGGCCGTTTGTGGCCCGACCTGAGTCAAAATCCGATCGTTGGACTCTTGGTG
CACCCCCCTTAGAGGAGGGATATGTGGTTCTGGTAGGAGACGAGAACCTAAAACAGTCC
CGCCTCCGTCTGAATTGGCTTCCGGTTGGGACCGAAGCCGCCGCGCTTGTCT
GCTGCAGCATCGTCTGTGTTGCTCTGTACTGTGTTCTGTATTGCTGTAAATA
TCGGCCCGGGCAGACTGTTACCACTCCCTTAAGTTGACCTTAGGTCACTGGAAAGATG
TCGAGCGGATCGCTACAACCAGTCGGTAGATGTCAAGAAAGAGACGTTGGGTTACCTCT
GCTCTGCAGAATGGCCAACCTTAACGTCGGATGGCCGCGAGACGGCACCTTAACCGAG
ACCTCATCACCCAGGTTAAGATCAAGGTCTTTCACCTGGCCCGCATGGACACCCAGACC
AGGTCCCCTACATCGTACCTGGGAAGCCTGGCTTTGACCCCCCTCCGGTCAAGC
CCTTGACACCCCTAACGCTCCGCTCCTCTCCATCCGCCCCGTCTCTCCCCCTTG
AACCTCCTCGTTGACCCCGCCTCGATCCTCCCTTATCCAGCCCTCACTCCTCT
GCGCCCCATATGCCATATGAGATCTTATGGGGCACCCCCGCCCTGTAAACTTCC
CTGACCCCTGACATGACAAGAGTTACTAACAGCCCTCTCTCCAAAGCTCACTACAGGCTC
TCTACTTAGTCCAGCACGAAGTCTGGAGACCTCTGGCGCAGCCTACCAAGAACAACTGG
ACCGACCGGTGGTACCTCACCCCTACCGAGTCGGCGACACAGTGTGGGTCCGCCGACACC
AGACTAAGAACCTAGAACCTCGCTGGAAAGGACCTTACACAGTCCTGCTGACCAACCCCA
CCGCCCTCAAAGTAGACGGCATCGCGCTTGGATACACGCCGCCACGTGAAGGCTGCCGA
CCCCGGGGTGGACCATCCTCTAGACTGCCGGATCTCGAGGGATCCACCACCATGGACCC
CCATTAAATTGGAATTCCCTGCAGCCCCGGGGATCCACTAGTTCTAGAGCGAATTAAATTCC

FIGURE 11A-2

GGTTATTTCCACCATATTGCCGTCTTGGCAATGTGAGGGCCCGAACCTGGCCCTG
TCTTCTGACGAGCATTCTAGGGTCTTCCCTCTGCCAAAGGAATGCAAGGTCTGT
TGAATGTCGTGAAGGAAGCAGTTCCTCTGGAAGCTTCTGAAGACAAACACGTCTGTAG
CGACCCCTTGAGGCAGCGGAACCCCCCACCTGGCAGACAGGTGCCCTGCGGCCAAAGC
CACGTGTATAAGATACACCTGCAAAGGCGGACAACCCAGTGCACGTTGTGAGTTGGA
TAGTTGTGAAAGAGTCAAATGGCTCTCTCAAGCGTATTCAACAAGGGCTGAAGGATG
CCCAGAAGGTACCCCATTGTATGGGATCTGATCTGGGCCCTCGGTGCACATGCTTACAT
GTGTTAGTCGAGGTTAAAAAAAGTCTAGGCCCCCGAACCAACGGGACGTGGTTTCCCT
TTGAAAAACACGATGATAATATGGGGATCCACCGGTGCCACCATGGTGAGCAAGGGCG
AGGAGCTGTCACCGGGTGGTGCCTACCTGGTCGAGCTGGACGGCACGTAACCGGC
ACAAGTTACGGTGTCCGGCAGGGCGAGGGCAGTGCACCTACGGCAAGCTGACCCCTGA
AGTTCATCTGCACCAACCGGAAGCTGCCGTGCCCTGGCCACCCCTCGTACCGACCCCTGA
CCTACGGCGTGCAGTGCTTCAGCCGTACCCGACCATGAAAGCAGCACGACTTCTCA
AGTCCGCCATGCCGAAGGCTACGTCCAGGAGGCACCATCTTCTCAAGGACGACGGCA
ACTACAAGACCGCGCGAGGTGAAGTTCGAGGGCGACACCCCTGGTAACCGCATCGAGC
TGAAGGGCATCGACTTCAAGGAGGACGGAACATCCTGGGGCACAAAGCTGGAGTACA
ACAACAGCCACAACGTCTATATCATGGCGACAAGCAGAAGAACGGCATCAAGGTGA
TCAAGATCCGCCACAACATCGAGGACGGCAGCGTGCAGCTGCCGACCACTACCAGCAGA
ACACCCCCATCGGCAGGGCCCCGTGCTGCCGACAACCAACTACCTGAGCACCCAGT
CCGCCCTGAGCAAAGACCCCAACGAGAACGGCGATCACATGGCCTGCTGGAGTTCGTGA
CCGCCGCCGGGATCACTCTGGCATGGACGAGCTGTACAAGTAAAGCAGGCCCTCGACGA
TAAAATAAAAGATTTATTAGTCTCCAGAAAAAGGGGGATGAAAGACCCACCTGTA
GGTTGGCAAGCTAGCTTAAGTAACGCCATTGCAAGGCATGGAAAAATACATAACTGA
GAATAGAGAAGTTCAAGGTCAAGGAGGACGGAACATGGAACAGCTGAATATGGGCCAA
GGATATCTGTGTAAGCAGTTCTGCCCGGCTCAGGGCCAAGAACAGATGGAACAGCTG
AATATGGGCCAAACAGGATATCTGTGTAAGCAGTTCTGCCCGGCTCAGGGCCAAGAA
CAGATGGTCCCCAGATGCGGCCAGCCCTCAGCAGTTCTAGAGAACCATCAGATTT
CAGGGTCCCCAAGGACCTGAAATGACCTGTGCCATTGAACTAACCAATCAGTT
CTTCTCGCTTCTGTTCGCGCTTCTGCTCCCCGAGCTCAATAAAAGAGCCCACAACCC
TCACTCGGGCGCCAGTCCGATTGACTGAGTCGCCGGTACCGTGTATCCAATAA
ACCCCTTGCACTGCATCCGACTTGTGGTCTCGCTGTTGGAGGGTCTCCTCTGA
GTGATTGACTACCCGTCAAGGGGGCTTTCATTCCGACTTGTGGTCTCGCTGCCATT
GAGGGTCTCCTCTGAGTGATTGACTACCCGTCAAGGGGGCTTCACTGAGCAGTAT
CAAATTAATTGGTTTTCTTAAGTATTACATTAAATGGCCATAGTTGCATTAAT
GAATCGGCCAACCGCGGGGAGAGGCAGGTTGCGTATTGGCGCTTCCGCTTCCTCGCT
CACTGACTCGCTCGCCTGGCTCGCTGCCGAGCGGTATCAGCTCACTAACAGG
GGTAATACGGTTATCCACAGAACAGGGATAACGCAGGAAAGAACATGTGAGCAAAGG
CCAGCAAAAGGCCAGGAACCGTAAAAGGCCGCGTTGCTGGCTTTCCATAGGCTCCG
CCCCCTGACGAGCATCACAAACATCGACGCTCAAGTCAGAGGTGGCGAACCCGACAGG
ACTATAAAGATACCAGGCCTTCCCCCTGGAAGCTCCCTCGTGCCTCTCTGTT
CCTGCCGCTTACCGGATACCTGTCGCCATTCTCCCTCGGAAGCGTGGCGCTTCTCA
TAGCTCACGCTGTAGGTATCTCAGTTCGGTAGGTCGTTGCTCCAAGCTGGCTGT
GCACGAACCCCCCGTTCAAGCCGACCGCTGCCATTCCGTAACTATCGTCTGAGTC
CAACCCGGTAAGACACGACTTATGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAG
AGCGAGGTATGTAGGCGGTGCTACAGAGTTCTGAAGTGGCTTAACACTACGGCTACAC
TAGAAGGACAGTATTGGTATCTCGCCTCTGCTGAAGCCAGTTACCTCGGAAAGAG
TGGTAGCTTGTACCGGAAACAAACCACCGCTGGTAGCGGTGTTTTGTTGCAA
GCAGCAGATTACCGCAGAAAAAGGATCTCAAGAACAGATCCTTGATCTTCTACGGG
GTCTGACGCTCAGTGGAACGAAAACACGTTAAGGGATTGGTACGAGATTATCAA
AAGGATCTCACCTAGATCCTTAAATTAAAAAGTAAAGTTGCCAAATCAATCTAAAG
TATATATGAGTAAACCTGGTCTGACAGTTACCAATGCTTAATCAGTGAGGCCACCTATCTC
AGCGATCTGTCTATTGCTCATCCATAGTTGCCGTGACTCCCCGCTGTAGATAACTAC
GATAACGGGAGGGCTTACCATCTGGCCCCAGTGCACGTTAAGGGATTGGTACGAGATTATCAA
ACCGGCTCCAGATTATCAGCAATAAACAGGCCAGCCGAAGGGCCGAGCGCAGAAGTGG

FIGURE 11A-3

TCCTGCAACTTATCCGCCTCCATCCAGTCTATTAATTGTTGCCGGGAAGCTAGAGTAAG
TAGTTGCCAGTTAATAGTTGCGCAACGTTGCTTGCATTGCTACAGGCATCGTGGTGTAC
ACGCTCGTCGTTGGTATGGCTTCATTCAAGCTCCGGTTCCCAACGATCAAGGCAGTTAC
ATGATCCCCCATGTTGTGCAAAAAAGCGGTTAGCTCCTTCGGTCCTCGATCGTTGTCAG
AAGTAAGTTGGCCGCAGTGTATCACTCATGGTTATGGCAGCACTGCATAATTCTCTAC
TGTCAATGCCATCCGTAAGATGCTTTCTGTGACTGGTGAGTACTCAACCAAGTCATTCTG
AGAATAGTGTATGCGCGACCGAGTTGCTCTGCCCCGGCGTCAACACGGGATAATACCGC
GCCACATAGCAGAACTTTAAAAGTGTCACTCATGGAAAAGTTCTCGGGCGAAAAGT
CTCAAGGATCTTACCGCTGTTGAGATCCAGTTGATGTAACCCACTCGTGACCCAACTG
ATCTTCAGCATCTTTACTTTACCAGCGTTCTGGGTGAGCAAAAACAGGAAGGCAAAA
TGCCGCAAAAAGGGAATAAGGGCGACACGGAAATGTTGAATACTCATACTCTCCTTT
TCAATATTATTGAAGCATTATCAGGGTTATTGTCTCATGAGCGGATACATATTGAATG
TATTTAGAAAAATAACAAATAGGGGTTCCCGCAGATTC

FIGURE 11B-1

1-845 CMV promoter/R/U5 5' LTR
1322 GAG ATG-ATC mutation
850-2100 extended □ region
2151-2865 GFP coding region
2866-2894 GGGSGGG linker
2895-2952 FMDV 2a cleavage sequence
2953-3004 BstX1/BstX1/HinD3/Hpa1/Sal1/Not1 polylinker
3052-3645 3' LTR
3652-5715 pGEM backbone (pUC origin, ampR)

ATCACGAGGCCCTTCGTCTCAAGAACAGCTTGCTCTAGGAGTTCTAAATACATC
CCAAACTCAAATATATAAGCATTTGACTTGTCTATGCCCTAGTTATTAAATAGTAATC
AATTACGGGGTCATTAGTTCATAGCCCCATATGGAGTCCCGTACATAACTTACGG
TAAATGGCCCGCCTGGCTGACCCCAACGACCCCCGCCATTGACGTCAATAATGACG
TATGTTCCCATACTAACGCCAATAGGGACTTCCATTGACGTCAATGGTGGAGTATT
ACGGTAAACTGCCACTTGGCAGTACATCAAGTGTATCATATGCCAAGTACGCCCTA
TTGACGTCAATGACGGTAAATGGCCCGCCTGGCATTATGCCAGTACATGACCTTATGG
GACTTCCCTACTTGGCAGTACATCTACGTATTAGTCATCGCTATTACCATGGTATGCG
GTTTTGGCAGTACATCAATGGCGTGGATAGCGGTTGACTCACGGGATTCCAAGTC
TCCACCCCATTGACGTCAATGGGAGTTGTTGGCACCAAAATCAACGGACTTCCA
AAATGTCGTAACAACCTCCGCCCCATTGACGCAAATGGCGGTAGGCATGTACGGTGGGA
GGTCTATATAAGCAGAGCTCAATAAAAGAGCCCACAACCCCTCACTCGGGCGCCAGTC
CTCCGATTGACTGAGTCGCCGGTACCCGTGTATCCAATAACCCCTTGCAGTTGCA
TCCGACTTGTGGTCTCGCTGTTCCCTGGGAGGGTCTCCTCTGAGTGATTGACTACCGT
CAGCGGGGGTCTTCATTGGGGCTCGTCCGGATCGGGAGACCCCTGCCAGGGACC
ACCGACCCACCACCGGGAGGTAAGCTGCCAGCAACTATCTGTCTGTCGATTGTC
TAGTGTCTATGACTGATTATGCGCCTGCGTCGGTACTAGTTAGCTAACTAGCTCTGT
ATCTGGCGGACCCGTGGAACTGACCGAGTCGGAACACCCGGCCGAAACCCCTGGGAG

FIGURE 11B-2

ACGTCCCAGGGACTTCGGGGCCGTTTGTGGCCCGACCTGAGTCCAAAATCCCGAT
CGTTTGACTCTTGGTGCACCCCCCTAGAGGAGGGATATGTGGTTCTGGTAGGAGA
CGAGAACCTAAAACAGTCCCGCCTCCGTCTGAATTGGCTTCGGTTGGGACCGAA
GCCGCGCCGCGCGTCTTGTCTGCTGCAGCATCGTCTGTGTTCTGTACTGTG
TTTCTGTATTGTCTGAAAATATCGGCCGGCCAGACTGTTACCACTCCCTTAAGTAA
GACCTTAGGTCACTGGAAAGATGTCGAGCGGATCGCTCACAAACAGTCGGTAGATGTCA
AGAAGAGACGTTGGTTACCTTCTGCTGCAGAATGGCCAACCTTAACGTCGGATGG
CCCGAGACGGCACCTTAACCGAGACCTCATCACCCAGGTTAAGATCAAGGTCTTTC
ACCTGGCCCGCATGGACACCCAGACCAGGTCCCACATCGTGACCTGGGAAGCCTTGG
CTTTGACCCCCCTCCCTGGGTCAAGCCCTTGTACACCCCTAACGCTCCGCCTCCTTT
CCTCCATCCGCCCGTCTCTCCCCCTTGAACCTCCTCGTTCGACCCCGCCTCGATCCCTC
CCTTATCCAGCCCTCACTCCTCTCTAGGCGCCCCATATGGCCATATGAGATCTTAT
ATGGGGCACCCCCGCCCCCTGTAAACTTCCCTGACCCCTGACATGACAAGAGTTACTAAC
AGCCCTCTCTCCAAGCTCACTTACAGGTCTCTACTTAGTCCAGCACGAAGTCTGGAG
ACCTCTGGCGGCAGCCTACCAAGAACAACTGGACCGACCGGTGGTACCTCACCCCTACC
GAGTCGGCGACACAGTGTGGTCCGCCGACACCAGACTAAGAACCTAGAACCTCGCTGG
AAAGGACCTTACACAGTCTGCTGACCACCCCCACCGCCCTAAAGTAGACGGCATCGC
AGCTTGGATAACACGCCGCCACGTGAAGGCTGCCGACCCGGGGTGGACCATCCTCTA
GAUTGCCGGATCTCGAGGGATCCACCATGGTGAAGCAAGGGCGAGGAGCTGTTACCGGG
GTGGTGCCTCATCCTGGTCAGGCTGGACGGCGACGTAAACGCCACAAGTTACGCGTGT
CGCGAGGGCGAGGGCGATGCCACCTACGGCAAGCTGACCCCTGAAGTTACGCGTGT
CCGGCAAGCTGCCGTGCCCTGCCACCCCTCGTGAACCACTACGGCGTGCAG
TGCTTCAGCCGCTACCCGACCACATGAAGCAGCACGACTTCTCAAGTCCGCATGCC
CGAAGGCTACGTCCAGGAGCGCACCATCTTCAAGGACGACGGCAACTACAAGACCC
GCGCCGAGGTGAAGTTGAGGGCGACACCCCTGGTGAACCGCATCGAGCTGAAGGGCATC
GACTTCAAGGAGGACGGCAACATCCTGGGCACAAGCTGGAGTACAACACTACAACAGCCA
CAACGTCTATATCATGCCGACAAGCAGAAGAACGGCATCAAGGTGAACCTCAAGATCC
GCCACAACATCGAGGACGGCAGCGTGCAGCTGCCGACCACTACCAGCAGAACACCCCC
ATCGGCACGGCCCGTGTGCTGCCGACAACCAACTACCTGAGCACCCAGTCCGCC
GAGCAAAGACCCCAACGAGAAGCGCGATCACATGGTCTGCTGGAGTTCGTACCGCC
CCGGGATCACTCTGGCATGGACGAGCTGTACAAGGAATTGGAGGTGGCAGCGGTGGC
GGTCAGCTGTTGAATTGGACCTTCTTAAACTTGGGGAGACGTCAGTCCAACCTGG
GCCACCACCATGGAAGCTCCATTAAATTGGTTAACGTCGACGCCGCGCTCGAC
GATAAAATAAAAGATTGTTAGTCTCCAGAAAAAGGGGGAAATGAAAGACCCACCT
GTAGGTTGGCAAGCTAGCTTAAGTAACGCCATTGGCAAGGCATGGAAAAATACATAA
CTGAGAATAGAGAAGTTCAGATCAAGGTCAAGGAAACAGATGGAACAGCTGAATATGGGCC
AAACAGGATATCTGTGTTAAGCAGTCCCTGCCCGCTCAGGGCCAAGAACAGATGGAA
CAGCTGAATATGGCCAACAGGATATCTGTGTTAAGCAGTCCCTGCCCGCTCAGGG
CCAAGAACAGATGGTCCCCAGATGCCGTCAGCCCTCAGCAGTCTAGAGAACCATCA
GATGTTCCAGGGTGCCCAAGGACCTGAAATGACCCCTGTCCTTATTTGAACCA
ATCAGTTCGCTCTCGTTCTGTCGCGCTTCTGCTCCCGAGCTCAATAAAAGAGC
CCACAACCCCTCACTCGGGCGCCAGTCCTCGATTGACTGAGTCGCCGGGTACCGT
GTATCCAATAAACCCCTTTGCAAGTGCAGTGCATCCGACTTGTGGTCTCGCTGTTCTGG
GGTCTCCTCTGAGTGAATGACTACCCGTCAAGCGGGGTCTTCATTTGACTACCGT
CTCGCTGCCTGGGAGGGTCTCCTCTGAGTGAATGACTACCCGTCAAGCGGGGTCTCA
CATGCAGCATGTATCAAATTAATTGGTTTTCTTAAGTATTACATTAAATGGC
CATAGTTGCATTAATGAATCGGCCAACCGCGGGAGAGGCGGGTTGCGTATTGGCGCT

FIGURE 11B-3

CTTCCGCTTCCTCGCTCACTGACTCGCTCGCTCGTCGGCTCGGGAGCGGTA
TCAGCTCACTCAAAGCGGTAATACGGTTATCCACAGAATCAGGGATAACGCAGGAAA
GAACATGTGAGCAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAGGCCGCTTGCTGG
CGTTTTCCATAGGCTCCGCCCTGACGAGCATCACAAAATCGACGCTCAAGTCAG
AGGTGGCGAAACCGACAGGACTATAAAGATAACCAGGCCTTCCCCCTGGAAGCTCCCT
CGTGCCTCTCCTGTTCCGACCCCTGCCGCTACCGGATACCTGTCCGCCTTCTCCCTT
CGGGAAAGCGTGGCGCTTCTCATAGCTCACGCTGTAGGTATCTCAGTCGGTAGGTC
GTTCGCTCCAAGCTGGCTGTGCACGAACCCCCGTTCAGCCGACCGCTGCGCCTT
ATCCGGTAACTATCGTCTGAGTCCAACCCGGTAAGACACGACTATGCCACTGGCAG
CAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTAGGCCTGCTACAGAGTTCTG
AAAGTGGTGGCTTAACACTACGGCTACACTAGAAGGACAGTATTGGTATCTGCCTCTGCT
GAAGCCAGTTACCTCGAAAAAGAGTTGGTAGCTCTGATCCGGAAACAAACCACCG
CTGGTAGCGGTGGTTTTGTTGCAAGCAGCAGATTACGCGCAGAAAAAAAGGATCT
CAAGAAGATCCTTGATCTTCTACGGGTCTGACGCTCAGTGGAACGAAAACACG
TTAAGGGATTTGGTCATGAGATTATCAAAAAGGATCTCACCTAGATCCTTTAAATT
AAAAATGAAGTTGCGCAAATCAATCTAAAGTATATATGAGTAAACTGGTCTGACAGT
TACCAATGCTTAATCAGTGAGGCACCTATCTCAGCGATCTGTCTATTCGTTCATCCAT
AGTTGCCCTGACTCCCCGTCGTAGATAACTACGATAACGGAGGGCTTACCATCTGGCC
CCAGTGCTGCAATGATAACCGCGAGACCCACGCTCACCGGCTCCAGATTATCAGCAATA
AACCAAGCCAGCCGAAGGGCGAGCGCAGAAGTGGTCTGCAACTTTATCCGCCTCCAT
CCAGTCTATTAATTGTTGCCATTGCTACAGGCATCGTGGTGTACGCTCGTCTGGTATGGCT
GCAACGTTGTTGCCATTGCTACAGGCATCGTGGTGTACGCTCGTCTGGTATGGCT
TCATTAGCTCCGGTCCCAACGATCAAGGCAGGTTACATGATCCCCATGTTGTGCAA
AAAAGCGGTTAGCTCCTCGGTCTCCGATCGTTGTACAGAAGTAAGTTGGCCGCAGTGT
TATCACTCATGGTTATGGCAGCACTGCATAATTCTTACTGTATGCCATCCGTAAGA
TGCTTTCTGTGACTGGTGAGTACTCAACCAAGTCATTCTGAGAATAGTGTATGCGGCG
ACCGAGTTGCTCTGCCGGCGTCAACACGGGATAATACCGGCCACATAGCAGAACTT
TAAAAGTGCTCATCTGGAAAAGCTTCTCGGGCGAAAACGATCTCAAGGATCTTACCG
CTGTTGAGATCCAGTTGATGTAACCCACTCGTGCACCCAACGATCTCAGCATCTT
TACTTCAACCAGCGTTCTGGGTGAGCAAAACAGGAAGGCAAAATGCCGAAAAAGG
GAATAAGGGCGACACGGAAATGTTGAATACTCATACTCTCCCTTTCAATATTATTGA
AGCATTATCAGGGTTATTGTCTCATGACATTAACCTATAAAATAGGCAGT

FIGURE 11C-1

1-845 CMV promoter/R/U5 5' LTR
1322 GAG ATG-ATC mutation
850-2100 extended 5' region
2146-2173 two BstX1 peptide cloning sites
2173-2214 EcoRI/ApaI/HpaI/NotI polylinker
2262-2855 3' LTR
2855-4901 pGEM backbone (pUC origin, ampR)

ATCACGAGGCCCTTCGTCTCAAGAACAGCTTGCTTCTAGGAGTTCTAATACATCCAAACTCAAAT
ATATAAAGCATTGACTTGTCTATGCCCTAGTTATTAAAGTAATCAATTACGGGTCAATTAGTTCATAG
CCATATATGGAGTTCCGCGTTACATAACTTACGGTAAATGCCCGCCTGGCTGACCGCCCAACGACCCCCG
CCCATTGACGTCAATAATGACGTATGTTCCCATAGTAACCCAATAGGGACTTTCCATTGACGTCAATGGG
TGGAGTATTACGGTAAACTGCCACTTGGCAGTACATCAAGTGTATCATATGCCAAGTACGCCCTATT
GACGTCAATGACGGTAAATGCCCGCCTGGCATTATGCCAGTACATGACCTTATGGGACTTTCTACTTG
GCAGTACATCTACGTATTAGTCATCGCTATTACCATGGTGTATGCCAGTACATCAATGGGCGTG
GATAGCGGTTTGACTCACGGGATTCCAAGTCTCCACCCATTGACGTCAATGGGAGTTGGCAC
CAAATCAACGGGACTTCAAAATGTCGAACAACCTCCGCCATTGACGCAAATGGGCGTAGGCATGT
ACGGTGGGAGGTCTATATAAGCAGAGCTCAATAAAAGAGCCCACAACCCCTCACTCGGGGCCAGTC
CGATTGACTGAGTCGCCGGTACCCGTGTATCCAATAACCCCTTGCAGTTGCATCCGACTTGTGGCT
CGCTGTTCTGGGAGGGTCTCTCTGAGTGTACTACCGTCAGGGGGCTTTCATTTGGGGCTC
GTCCGGGATCGGGAGACCCCTGCCAGGGACCACCGACCCACCACGGGAGGTAAGCTGGCAGCAACTTA
TCTGTGCTGCTGATTGCTAGTGTCTATGACTGATTTATGCCCTGCGTGTACTAGTTAGCTA
AGCTCTGTATCTGGCGGACCCGTGGTGAAGTGAAGCTCGGAACACCCGGCGCAACCCGGAGACGT
CCCAGGGACTCGGGGGCCCTTTGTGGCCGACCTGAGTCCAAAATCCGATCGTTGGACTTTG
GTGCACCCCCCTAGAGGAGGGATATGTGGTCTGGTAGGAGACGAGAACCTAAAACAGTCCCGC
TCTGAATTTTGCTTCCGTTGGGACCGAAGCCGCGCCGCGTCTGTCTGCTGCAGCATCGTTGTG
TTGCTCTGTCTGACTGTGTTCTGTATTGTCTGAAAATATCGGCCCCGGCAGACTGT
TAAGTTGACCTAGGTCACTGAAAGATGTCGAGCGGATCGCTCACAACCAGTCGGTAGATGTC
AGACGTTGGTTACCTCTGCTCTGCGAAATGCCAACCTTAACGTCGGATGGCGAGACGGCACCT
TAACCGAGACCTCATCACCAGGTTAAGATCAAGGTCTTCAACCTGCCGATGGACACCCAG
TCCCTACATCGTGAACCTGGGAAGCCTGGCTTTGACCCCCCTCCCTGGTCAAGCCCTTG
AAGCCTCCGCTCCTCTCCATCCGCCGCTCTCCCCCTTGACCTCGTCTGACCCCCGCC
ATCCTCCCTTATCCAGCCCTACTCTCTAGGCGCCCCCATATGCCATATGAGATCTTATATGGG
CACCCCCGCCCTGTAAACTTCCCTGACCTGACATGACAAGAGTACTAACAGCCCTCTCCAAGCT
CACTTACAGGCTCTACTTAGTCCAGCAGAAGTCTGGAGACCTCTGGCGGAGCCTACCAAGAAC
GGACCGACCGGTGGTACCTACCCCTTACGAGTCGGCGACACAGTGTGGTCCGGCAGAC
ACCTAGAACCTCGCTGGAAAGGACCTTACACAGTCCTGCTGACCCACCCGCC
ATCGCAGCTGGATACACGCCGCCACGTGAAGGCTGCCGACCCGGGGTGGACC
GGATCTCGAGGGATCCACCAACATGGACCCCCATTAAATTGGAA
ACGCGGCCGCCGTGACGATAAAATAAAAGATT
CACCTGTAGTTGGCAAGCTAGCTTAAGTAACGCCATTGCAAGGCATGGAAA
TAGAGAAGTTAGCTAGATCAAGGTCAGGAACAGATGGAACAGCTGA
AGCAGTTCTGCCCGCTCAGGGCCAAGAACAGATGGAACAGCTGA
GGTAAGCAGTTCTGCCCGCTCAGGGCCAAGAACAGATGGTCCCC
TTCTAGAGAACCATCAGATGTTCCAGGGTGCCCCAAGGAC
ACCAATCAGTTCCCTCTGCTCTGCTGCCGCTTCTGCT
CCTCACTCGGGCGCCAGTCCCGATTGACTGAGTC
GCCGGGTACCGTGATCCAATAACCCCTTTG

FIGURE 11C-2

CAGTTGCATCCGACTTGTGGCTCGCTGTTCTGGGAGGGCTCCTCTGAGTGATTGACTACCCGTCA
GGGGGTCTTCATTTCCGACTTGTGGCTCGCTGCCCTGGGAGGGCTCCTCTGAGTGATTGACTACCCGT
CAGCGGGGTCTTCACATGCAGCATGTATCAAATTAATTGGTTTTCTTAAGTATTACATTAAT
GCCATAGTTGCATTAATGAATCGGCCAACCGCGGGGAGAGGCGGTTGCGTATTGGCGCTCTCCGCTT
CCTCGCTCACTGACTCGCTCGCCTCGGCTGCGCTGCGAGCGGTATCAGCTCACTCAAAGGCCAGTA
ATACGGTTATCCACAGAATCAGGGATAACCGCAGGAAAGAACATGTGAGCAGAAAAGGCCAGCAAAGGCCAG
GAACCGTAAAAGGCCCGTGTGGCTTTCATAGGCTCCGCCCCCTGACGAGCATCACAAAATC
GACGCTCAAGTCAGAGGTGGCGAACCCGACAGGACTATAAAGATACCAAGGCGTTCCCCCTGGAAGCTCC
CTCGTGCCTCTCTGTTCCGACCCCTGGCCTAACGGATACCTGTCCGCTTCTCCCTCGGAAGCGT
GGCGTTCTCATAGCTCACGCTGTAGGTATCTCAGTTGGCTAGGCTCGCTCCAAAGCTGGCTGTG
TGCACGAACCCCCCGTTCAGCCGACCGCTGCGCCTTATCCGTAACTATCGTCTGAGTCCAACCCGGTA
AGACACGACTTATGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTAGGCGGTGC
TACAGAGTTCTGAAGTGGTGGCTAACACGGCTACACTAGAAGGACAGTATTGGTATCTCGCTCTGC
TGAAGCCAGTTACCTCGGAAAAGAGTTGGTAGCTCTTGATCCGAAACAAACCCACCGCTGGTAGCGGT
GGTTTTTTGTTGCAAGCAGCAGATTACCGCAGAAAAAAAGGATCTCAAGAAGATCCTTGATTTTC
TACGGGTCTGACGCTCAGTGGAACGAAAACACGGTTAACGGATTGGTATGAGATTATCAAAAAGGA
TCTTCACCTAGATCCTTTAAATTAAAAATGAAGTTGCGCAAATCAATCTAAAGTATATGAGTAAACT
TGGTCTGACAGTTACCAATGCTTAATCAGTGGCACCTATCTCAGCGATCTGTCTATTGCTCATCCAT
AGTTGCTGACTCCCCGTCGTAGATAACTACGATACGGGAGGGCTTACCATCTGGCCCCAGTGTGCAA
TGATACCGCAGACCCACGCTCACCGGCTCCAGATTATCAGCAATAAACAGCCAGCCGGAAAGGGCCGAG
CGCAGAAGTGGTCTGCAACTTATCCGCTCCATCCAGTCTATTAAATTGTTGCCGGAAAGCTAGAGTAAG
TAGTTGCCAGTTAATAGTTGCGCAACGTTGCTTGCCTTGCTACAGGCATCGTGGTGTACGCTCGTGT
TTGGTATGGCTTCATTGCTCCGGTCCACGATCAAGGCGAGTTACATGATCCCCATGTTGCAA
AAAGCGTTAGCTCCTTCGGCCTCCGATCGTTGTCAGAAGTAAGTTGGCCGAGTGTATCACTCATGGT
TATGGCAGCACTGCATAATTCTCTTACTGTCATGCCATCCGTAAGATGCTTTCTGTGACTGGTAGTACT
CAACCAAGTCATTGAGAATAGTGTATGCGGCCACCGAGTTGCTCTGCCCCGGTCAACACGGGATAAT
ACCGGCCACATAGCAGAACCTTAAAGTGTCTCATGGAAAACGTTCTCGGGCGAAAACCTCTCAAG
GATCTTACCGCTGTTGAGATCCAGTTGATGTAACCCACTCGTCAGCCAACTGATCTCAGCATCTTTA
CTTTCACCAGCGTTCTGGGTGAGCAAAACAGGAAGGCAAAATGCCCAAAAGGAAATAAGGGCGACA
CGGAAATGTTGAATACTCATACTCTCCTTTCAATATTATTGAAGCATTATCAGGGTTATTGTCTCAT
GACATTAACCTATAAAAATAGGCAGT

FIG. 12 A

(1) C12ScFas Survival construct

C12ScFas: epsilon-cFas(CD95)-Ires-Hygro-BGH PolyA put into C12s vector backwards so that no leaky transcription happens through the cmv promoter. .

F16 12 B

TTTTTACCAAGGTTGGCATGGTGCAGACGAAATGGGCCTCCTGATATAATCCTCTGAGCAGTTTATCAGTTCATG
AACCCGCCTCCTCAGCTTAAACTCTGGAGATGCTATTAGTACCTTGAGTATGAACCTTAACGTGAGCCAGCAAGCA
CCAGAGGCAGGACAGCCCAGATCCACACCATgGTGGCTTACCAACAGTACCGGAATGCCAAGCTGCGGCCCTTAAGA
GCTGTAATTGAACCTGGGAGTGGACACCTGTGGAGAGAAAGCAAAGTGGATGTCAGTAAGACCAATAGGTGCCATCAG
AAACGCAAGAGTCTCTGTCTGACAAGCCCAGTTCTATTGGTCTCCTAAACCTGTCTGTAAACCTTGATACCTAC
CTGCCCAGTGCCCTCACGACCAACTTctgcaggattcctggacagctcccgatgtcgtacgtaccgtggtttttct
gtccggggcagtggagccctggtagggggagctgcctcagtgcctcgtttcagctaaaatgggggggggggggggggggg
cccggggccctggaaaggcc
ttcactaccatgcacaccggagtgtGggggaggttcttcgtcggcccaacCccagggccctgccttaggtcccc
ggactctCactttgcgcgtgcgtggcttgcgtggccctcggccctcggccctcggccctcggccctcggccctcggcc
ggacttgggcatgcgcgtgcgtggccctcggccctcggccctcggccctcggccctcggccctcggccctcggcc
cgacttgggcatgcgcgtgcgtggccctcggccctcggccctcggccctcggccctcggccctcggccctcggcc
ctgtgacccgcttggagctggcaccctgagttggccacacCTTGTACTCACTCCCAGGTCACTGTCCtcacGCGGCC
GCTCGAcgatAAAATAAAAGATTTATTTAGTCTCCAGAAAAAGGGGGGAATGAAAGACCCACCTGTAGGTTGGCAAG
ctagcTTAAGTAACCCATTGCAAGGCATGGAAAAATACATAACTGAGAATAGAGAAGTTCAAGGTCGGAACAG
ATGGAACAGGAATAAAAGAGCCCACAACCCCTCACTCGGGCGCCAGTCCTCCGATGACTGAGTCGCCGGTACCCG
TGTATCCAATAACCCCTTGCAGTTGCATCCGACTTGTGGTCTCGCTGTTCTGGAGGGTCTCCTGTAGTATTGA
CTACCCGTAGCGGGGGTCTTCatgcacGCATGTATCAAAATTAAATTGGTTTTCTTAAGTATTACATTAAAT
GGCCATagttcGTAATCATGGTCATAGCTGTTCCCTGTGAAATTGTTATCCGTCACAATTCCACACAATACGAG
CCGGAAGCATAAAAGTGTAAAGCCTGGGTGCCTAATGAGTGAGCTAACTCACATTAAATTGCGTTGCGCTACTGCCGCT
TTCCAGTCGGAAACCTGCGCCAGCTGCATTAATGAATCGCCAACCGCGGGAGAGGCGGTTGCGTATTGGGCG
CTCTCGCTCCCTCGCTACTGACTCGCTGCGCTCGTCGTCGCGAGCGGTATCAGCTCACTCAAAGGCGG
TAATACGGTTATCCACAGAATCAGGGATAACGCAGGAAAGAACATGTGAGCAGCAGGCCAGGAAACCGT
AAAAAGGCCGCGTTGCTGGCTTTCCATAGGCTCCGCCCCCTGACGAGCATCACAAAATCGACGCTCAAGTCAGAG
GTGGCGAAACCGACAGGACTATAAGATACCAAGCGTTCCCTGGAAAGCGTGGCGCTTCATAGCTCACGCTGTAGGTATCTC
TGCGCTTACCGGATACCTGTCGCCCTTCTCCCTCGGAAGCGTGGCGCTTCAGCCGACCGCTGCGCTTATCCGG
TAACTATCGCTTGAGTCCAACCCGTAAGACACGACTTATGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAG
CGAGGTATGAGGCGGTGCTACAGAGTTCTGAAGTGGTGGCTAACTACGGCTACACTAGAAGGACAGTATTGGTATC
TGCCTCTGCTGAAGCCAGTTACCTCGGAAAAGAGTTGGTAGCTCTGATCCGCAAACAAACCACCGCTGGTAGCGG
TGGTTTTTTGTTGCAAGCAGCAGATTACGCGCAGAAAAAAAGGATCTCAAGAAGATCTTGTACAGTAAACTACGCT
CTGACGCTCAGTGGACGAAAACCTCACGTTAAGGGATTGGTCTGAGATTATCAAAGGATCTCACCTAGATCCTT
TTAAATTAAAAATGAAGTTGCGCAAATCAATCTAAAGTATATGAGTAAACTTGTGACAGTTACCAATGCTTAAT
CAGTGAGGCACCTATCTCAGCGATCTGTCTATTGCTTCCATAGTGCCTGACTCCCCGTCGTAGATAACTACGA
TACGGGAGGGCTTACCATCTGGCCCCAGTGCCTGCAATGATAACCGCAGACCCACGCTCACCGCTCCAGATTATCAGCA
ATAAACCCAGCCAGCCGAAGGGCCGAGCGCAGAAGTGGTCTGCAACTTATCCGCTCCATCCAGTCTATTAAATTGTTG
CCGGGAAGCTAGAGTAAGTAGTTCGCCAGTTAATAGTTGCGCAACGTTGCTGCAAGGCTACAGGATCGTGGTCAC
GCTCGTGTGTTGGTATGGCTTCACTCAGCTCCGGTCCAAAGGATCAAGGCGAGTTACATGATCCCCATGTTGCAAA
AAAGCGGTTAGCTCCTTCGGCCTCCGATGTTGTCAGAAGTAAGTTGCCAGTGTATCACTCATGGTATGGCAGC

FIG 12C

ACTGCATAATTCTCTTACTGTCATGCCATCCGTAAGATGCTTCTGTGACTGGTGagactcaaccaagtcattctgag
aatagtgtatcgccgacccgagttgtcttgcggcgtaacacacggataataccgcgccacatagcagaactttaaaa
gtgctcatcattggaaaacgttcttcggggcgaaaactctcaaggatcttaccgctgttagatccagtcgtatgtacc
cactcgtgcacccaactgtatcttcagcatctttactttcaccagcgttctgggtgagcaaaaacaggaaggcaaaatg
ccgcaaaaaggaaataagggcgacacggaaatgttgaatactcatactcttcctttcaatattatttgaagcatttat
cagggttattgtctcatgacattaacctataaaaataggcgt

ପାଇଁ ପାଇଁ ପାଇଁ ପାଇଁ ପାଇଁ ପାଇଁ ପାଇଁ ପାଇଁ ପାଇଁ

FIG 13A

(2) Ahhhh: Survival construct

2.) Ahhhh: epsilon-cFas' (CD8 or mLyt2)-Ires-Hygro-BGHpolyA also in C12s backwards

Fl- 13 B

tcaacatcagataaatttattgccactgtttcaggatthaaggttgagattcatgagaacctgggtttcccttcgt
cttctgcattttctgtactccttcttcacccaaacaatttagtggattggaaaagaagacaagccacc
ccaaaccgggttccggcccccactgagccacggggccgacaatctctggctctgggctgagatgtccggtaggg
tgcacaggtgaggagttcgacactggcttgtagtagatgtacttctgaaggactggcacgacagaactgaa
gtacatcaccgagttcgatgactgagcagaaatagtagccttcgtacttctgtcacttgcggtaggtgagaacgt
acttattattctgtggccatggcagaaaacatgtggccacgatcttcgtccacgttctgtccacgttatctgttgggat
gaagccatatacacaacgaagggtggctggggagttggagctggagttctggaaagagccaaagagcatcttgcgaaac
ggaccccaacactcacataccaggtccacccactgaccaagttcggcgtccatcttggaaagattcggagttcgg
gtgcctgtggcttagcttccactccccaggataatcgactcaccagcagcagggtagcgcacagaaacgggt
aacggtgaggccatgGTGGCTTTACCAACAGTACCGGAATGCCAAGCTGCGGCCGCTTAAGAGCTGTAATTGAACCTGG
GAGTGGACACCTGTGGAGAGAAAGGCAAAGTGGATGTCAGTAAGACCAATAGGTGCCTATCAGAAACGCAAGAGTCTTCT
CTGTCGACAAGCCCAGTTCTATTGGTCTCCTAACCTCTTGTAACTTACCTGCCAGTGCCTCACG
ACCAACTTctgcaggaattcctggacagctccagatgtactcgttaaccgtgggttatttctgtgcggggcagtggagc
ctggtagggggagctctgcctcagtcttcagctaaaatgggtggaaacccCaggaggccggccggccctggaa
gttccctttctctgttcttggagtcgatttagcaacacgccccacCcaggggccctgcctgtggactctCactcttgac
ccgagtgtGggggaggttctcttcgtggccacgtcggccaaacttgggtggctgtggaaaggagagggtactgggcacgc
gcacgtcgtggcttggccatgtggccatgtggccatgtggccatgtggccatgtggccatgtggccatgtggccatgtgg
gcctcgccccGccgtgtccagatgtgtgtcaggccctctgtggcccgccctgtggccatgtggccatgtggccatgtgg
ctggcaccctgtgtggccacatgt
AAGATTTATTTAGTCTCCAGAAAAAGGGGGAAATGAAAGACCCACCTGTAGGTTGGCAAGctagcTTAAGTAACCCA
TTTGCAAGGCATGGAAAAATACATAACTGAGAATAGAGAAGTCAGATCAAGGTCGAACAGATGGAACAGGAATAAA
AGAGCCCACACCCCTCACTCGGGGCCAGTCCTCCGATTGACTGAGTCGCCGGTACCGTGTATCCAATAACCC
CTTGCAGTTGCATCCGACTTGTGGCTCGCTGTTCTGGAGGGTCTCTGTAGTGATTGACTACCCGTCAAGCGGGGG
TCTTTCAcatgcaGCATGTATCAAATTAATTGGTTTTCTTAAGTATTTACATTAAATGCCATagttcGTAAT
CATGGTCATAGCTGTTCCGTGTGAAATTGTTATCCGCTACAATTCCACACAATACGAGCCGAAGCATAAGTGT
AAAGCCTGGGTGCTAATGAGTGGCTAACTCACATTAAATGCGTTGCCCTACTGCCGTTCCAGTCGGAAACCT
GTCGTGCCAGCTGCATTAATGAATCGCCAACGCCGGGGAGAGGCGGTTGCGTATTGGCGCTCTCCGCTTCG
TCACTGACTCGCTCGCTCGTCGGCTCGCTGCCGAGCGGTATCAGCTCACTCAAAGCGGTAAACGGTTATCCACA
GAATCAGGGATAACGCAAGGAAAGAACATGTGAGCAAAGGCCAGCAAAGGCCAGGAACCGTAAAAGGCCGTTGCT
GGCGTTTTCCATAGGCTCCGCCCCCTGACGAGCATCACAAAATCGACGCTCAAGTCAGAGGTGGCAAACCGACAG
GACTATAAAGATAACCAGGGTTTCCCTGGAAAGCTCCCTGTGCGCTCTCTGTGTTCCGACCCGCCCTACCGGATAC
CTGTCGCCCTTCTCCCTCGGGAAAGCGTGGCGTTCTCATAGCTCACGCTGTAGGTATCTCAGTTGGTAGGTG
TCGCTCCAAGCTGGCTGTGACGAACCCCCCTTCAGCCGACCGCTGCCCTATCCGTAACATCGTCTTGAGT
CCAACCCGGTAAGACACGACTTATGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGAGGCGGT
GCTACAGAGTTCTGAAGTGGTGGCTAACTACGGCTACACTAGAAGGACAGTATTGGTATCTGCGCTCTGCTGAAGCC
AGTTACCTCGAAAAAGAGTTGGTAGCTCTTGATCCGGCAAACAAACCCACCGCTGGTAGCGGTGGTTTTGTTG
AGCAGCAGATTACGCGCAGAAAAAGGATCTCAAGAAGATCCTTGATCTTCTACGGGTCTGACGCTCAGTGGAAAC
GAAAACTCACGTTAAGGGATTTGGTATGAGATTATCAAAAAGGATCTCACCTAGATCCTTAAATTAAAAATGAAG
TTTGCCTAACATCTAAAGTATATGAGTAAACTTGGTCTGACAGTTACCAATGCTTAATCAGTGGAGGACCTATCT
CAGCGATCTGCTTATTCGTTCATCCATAGTTGCTGACTCCCGTGTAGATAACTACGATAACGGAGGGCTTACCA
TCTGGCCCCAGTGCTGCAATGATAACCGCAGACCCACGCTCACGGCTCCAGATTTATCAGCAATAACCCAGCCAGCCGG
AAGGGCCGAGCCAGAAGTGGTGGCTGCAACTTATCCGCTCATCCAGTCTATTAAATTGTTGCCGGAAAGCTAGAGTAA
GTAGTTGCCAGTTAATAGTTGCGCAACGTTGCTACAGGATCGTGGTGTACGCTCGTGTGTTGGTATG

FIG 13C

GCTTCATTCAGCTCCGGTCCAAACGATCAAGGCGAGTTACATGATCCCCATGTTGTGCAAAAAAGCGGTTAGCTCCTT
CGGTCCCTCCGATCGTGTAGAAGTAAGTTGGCCGCAGTGTATCACTCATGGTTATGGCAGCAGTCATAATTCTCTTA
CTGPCATGCCATCCGTAAGATGCTTTCTGTGACTGGT~~Gagtactcaaccaagtcattctgagaatagtgtatgcggcga~~
~~ccgagttgctcttgcggcgtaaacacggataataccgcgcacatagcagaacttaaaagtgcgtcatcattggaaa~~
~~acgttcttcggggcgaaaactctcaaggatcttaccgcgttggatccagttcgatgtaaacccactcgtgcacccaaact~~
~~gatcttcagcatctttactttcaccagcggttctgggtgagcaaaaacaggaaggcaaaatgcccggaaaaaggaaata~~
~~agggcgacacgaaatgttgaataactcataactcttcctttcaatattattgaagcatttatcagggttattgtctcat~~
~~gacattaacctataaaaataggcgt~~